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APPLICATION NO.	FILING DATE	FIRST NAMED	INVENTOR		ATTORNEY DOCKET NO.	
09/125,841	01/19/99	OLSEN		R	CIR20013	
Г		1144070645	\neg		EXAMINER	
MUELLER & SMITH 7700 RIVERS EDGE DRIVE COLUMBUS OH 43235		HM12/0615	·	SCHWAD	SCHWADRON, R	
				ART UNIT	PAPER NUMBER	
		• •		1644	8	
				DATE MAILED:	06/15/00	

Please find below and/or attached an Office communication concerning this application or proceeding.

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SERIAL NUMBER	FILING DATE	FILING DATE FIRST NAMED APPLICANT	

EXA	MINER
ART UNIT	PAPER NUMBER

DATE MAILED:

Please find below a communication from the EXAMINER in charge of this application

Commissioner of Patents

The communication filed on 5/2/2000 is not fully responsive to the communication mailed 4/12/2000 for the reason(s) set forth on the attached Notice to Comply With the Sequence Rules or CRF Diskette Problem Report.

Since the response appears to be <u>bona fide</u>, but through an apparent oversight or inadvertence failed to provide a complete response, applicant is given **ONE** (1) **MONTH or THIRTY (30) DAYS** from the mailing date of this notice, whichever is longer, within which to supply the omission or correction in order to avoid abandonment. EXTENSIONS OF THIS TIME PERIOD MAY BE GRANTED UNDER 37 CFR 1.136(a).

Any inquiry concerning this communication should be directed to Examiner Ron Schwadron, Ph.D., Art Unit 1644, whose telephone number is 703-308-4680.

Any inquiry of a general nature or relating to the status of this application should be directed to the Technology Center receptionist whose telephone number is (703) 308-0196.

Rsh

RONALD B. SCHWADRON
PRIMARY EXAMINER
GROUP 1899- 1 600

Ron Schwadron, Ph.D. Art Unit 1644

Application No.: 09/125841

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

K	 This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
	This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
	3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
A	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
	5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
	6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
	7. Other:
Δn	plicant Must Provide:
X	An <u>initial</u> or substitute computer readable form (CRF) copy of the "Sequence Listing".
K	An <u>initial</u> or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
X	A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).
Fo	r questions regarding compliance to these requirements, please contact:

For questions regarding comp

For Rules Interpretation, call (703) 308-4216

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Sequence Rubs

Survey Llesse MANDATORY

numerie identifiers

and nos <212> DNA <213> HIV-1 primer <400> 1 ataatccacc tatcccagta ggagaaat 28 <210> 2 **Does Not Comply** Corrected Diskette Needed <211> 28 <212> DNA <213> HIV-1 primer <400> 2 tttggtcctt gtcttatgtc cagaatgc 28 <210> 3 <211> 22 <212> DNA <213> GADPH cDNA <400> 3 ctactggcgc tgccaaggct gt <210> 4 <211> 22 <212> DNA

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